GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

OM nucleic - nucleic search, using sw model

December 6, 2002, 22:41:46; Search time 2691.5 Seconds (without alignments) 16489.608 Million cell updates/sec Run on:

US-10-025-514-7 1525 Title: Perfect score:

1 tctagaccatgtctggaaag........ccaactcagaagtagtcgac 1525 Sequence:

Scoring table:

2054640 seqs, 14551402878 residues IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Searched:

4109280 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

GenEmb1:*

9b_ba:*
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em_htg_mam:*
em_htg_mam:*
em_htg_mam:*
em_htg_vr:*
em_sy:*
em_htgo_htm:*

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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ALIGNMENTS

1308 bp Sequence 6 from patent US 6127145. AR111412 AR111412.1 GI:12828260 Unknown. RESULT 1
AR111412
LOCUS
DEFINITION
ACCESSION
VERSION
KEYWORDS SOURCE ORGANISM

PAT 14-FEB-2001

linear

DNA

JM Unknown.
Unclassified.
Location/Qualifiers

Moclassified.
Location/Qualifiers

REFERENCE AUTHORS TITLE JOURNAL FEATURES

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                                                                              Length 1308;
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Pred. No. 2.1e-125;
0; Mismatches 346;
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                                1076 CGTGACGGAGGAGGCCCCCTGAAGCTCTCCAAGGCCGTGCACAAGGCGGTGCTCCACGAT 1135
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1355 TGATGAAAAGGGTACCGAGGCCGCCGCGCTATGTTCCTGGAAGCTATTCCAATGAGCAT
                                                                                                                                   1415 TCCACCAGAAGTTAAATTTAATAAACCATTCGTTTTTCTGATGATCGAGCAGAACACTAA
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Stalliff,T.D. and Rodriguez,R.L.
Production of alpha.sub.1 -antitrypsin in plants
Patent: US 6127145-A 3 03-OCT-2000;
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Sequence 3 from patent US 6127145.
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396 c 334 q
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Best Local Similarity 70.83
Matches 837; Conservative
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704 TITAAAATTGGTTGACAAATTCCTAGAAGACGTCAAGAACTATATCATAGTGAGGCTTT 763
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/db_xref="taxon:9606"
/map="14q32.1"
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/db_xref="G1:177827"
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alpha-1-antitrypsin.
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PEDGEKLGHLENELTHDIITKFLENEDRRSASLHLPKLSITGTYDLKSVLGQLGITKV
FSNGADLSGVTEEAFLLSKAVKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNKP
FVFLMIEQNTKSPLEMGKVVNPTQK"
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//translation="MPSSVSWGILLAGLCCLVPVSLAEDPQGDAAQKTDTSHHDQDHP
TFNKITPNLAEFAFSLYRQLAHQSNSTNIFFSPVSIATAFAMLSLGTKADTHDEILEG
INFNLTEIPEAQIHEGFQELLRTLNQPDSGLQLTTGNGLFLSEGLKLVDFFLEDVKL
[2] sites; alpha-1-antitrypsin deficiency disease mutations. The Stanite Falls (a one base pair delection), Bellingham (a substitution) and Mattawa (a one base pair insertion) mutations all result in premature transcription termination.
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Pred. No. 5.8e-85;
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//note='amino acid 358-Val'
replace(1190. 1193, GGG')
//note='amino acid 358-Gly'
replace(1190. 1193, AGA')
//note='amino acid 358-Erg'
replace(1190. 1193, AGA')
//note='amino acid 358-Erg'
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Magazetsuto, W. 1. and Guren, H.K.
REGION SPECIFIC MUTATION INDUCTION IN ALPHA-1-ANTI-TRYPSIN Patent: JP 1986012289-A 1 20-JAN-1986; CHIMO JIENETEITSUKUSU INC
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                                                                                            838 AGATCGTTGACCTAGTTAAAGAATTAGATCGTGATACCGTCTTCGCACTAGTTAACTATA 897
                                                                                                            Glycolytic promotersfor regulated protein expression: protease inhibitor
                                                                 561 GGGACACCGAAGAGGCCAAGAACAGATCAACGATTACGTGGAGAAGGGTACTCAAGGGA
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1547 - 16th Ave. East; Seattle, WA
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1 (bases 1 to 1312)
Kawasaki,G.H.
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                                          Length 1312;
                                         Score 436.4; DB 6; Length
Pred. No. 1.1e-83;
0; Mismatches 491; Indels
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.nhibitor, clade A (alpha-1 antiproteinase, antitrypsin),
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/db_xref="G1:15080499"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BC011991
Homo sapiens, Similar to serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1, clone MGC:9222 IMAGE:3859644, mRNA, complete cds.
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Contact: villalon@bcm.tmc.edu.
Villalon, D.K., Luna, R.A., Hale, S.M., Hulyk, S., Lu, X., Garcia,
A.M., Holloway, M., Telford, B, Hodgson, A., Bouck, J., Yu, W.,
Muzny, D.M., Glbbs, R.A.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1584)
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              1258 ITACCAAAGTITITICIAACGGIGCCGAITIGAGIGGIGTIACIGAAGAAGCICCAITAA 1317
                                                        1318 AATTGAGTAAAGCTGTTCACAAAGCCGTCTTAACTATTGATGAAAAGGGTACCGAGGCCG 1377
                                                                                                                                                                                                                   1082 ACCICICCAAGGCCGIGCAIAAGGCIGIGCIGACCAICGACGAGAAAGGGACIGAAGCIG 1141
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/product="Similar to serine (or cysteine) proteinase
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/clone_lib="NIH_MGC_66"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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Cancer gene determination and therapeutic screening using signature
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/Organism="Homo saplens"
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838 AGAICGIIGACCIAGIIAAAGAAITAGAICGIGAIACCGICTICGCACIAGIIAACIANA 897
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Pred. No. 5.6e-83;
0; Mismatches 493; Indels 0;
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90 bp upstream of DdeI site.
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Best Local Similarity 59.7%;
Matches 729; Conservative
/citation=[1]
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/gene="PI"
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PTHRITTPRAFFOLYGLAHQSTRINFFSPVSTATAFAMLSLGTKADTHDELLE
GLURNLTERPEAGLHCFFGELLRTLMQPDSGLQLTTGRWGLFLSBGTKLVDKFLEDVKK
LYHSEAFTVNFGDTEEAKKQINDXVEKGTGGKIVDLVKELDRDTVFALVNYIFFRGKW
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FLDDEGKLOHLENECTHDITTKFLENEDRRSASLHLPKLSITGTYDLKSVLGQLGITK
VFSNGADLSGVTEEAPLKLSKAVHKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNK
PFVFLMIEQNTKSPLEMGKVVNPTQK"
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                            PRI 01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Revised sequence of full-length complementary DNA coding for human
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1231)
2011en, A., Herzoag, A., Cravador, A., Herion, P., Chuchana, P., Vander Straten, A., Loriau, R., Jacobb, P. and van Elsen, A. Cloning and expression in Escherichia coli of full-length DNA 2 (4), 255-264 (1983)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Courtney, M., Buchwalder, A., Tessier, L.H., Jaye, M., Benavente, A., Balland, A., Kohli, V., Lathe, R., Tolstoshev, P. and Lecocq, J.P. High-level production of biologically active human alpha l-antitrypsin in Escherichia coli a. Froc. Natl. Acad. Sci. U.S.A. 81 (3), 669-673 (1984) 84144765
                                                                                                                       alpha-1-antitrypsin; antiprotease; antitrypsin.
Human liver, cDNA to mRNA, clones pULB1523 [1] and pTG603 [3]
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                                Human alpha-1-antitrypsin mRNA, complete cds.
K01396
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/db_xref="taxon:9606"
/map="14q32.1"
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TTTTTTCAAGGGTAAGTGGGAACGTCCTTTCGAGGTTAAAGATACTGAAGAGGAAGATT	TACCGAACTGTCCATTACTGGAACCTATGATCTGAAGAGCGTCCTGGGTCAACTGGGCA TTACCAAAGTTTTTTCTAACGTGCCGATTTGAGTGGTGTTACTGAAGAGCTCCATGA	CIGGGGCCAIGITTTAGAGGCCAIACCCAIGTCTATCCCCCCCGAGGTCAAGTTCAACA 1193 AACCATTGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCATTGTTTATGGGTAAGG 1497	AX335336 N Sequence 5847 from Patent W00194629. AX335338 AX335338 AX335338.1 GI:18126057 human. M Homo saplens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 Tong, P.E., Augustuas, M., Carter, K.C., Ebner, R., Endress, G., Tong, P.E., Augustuas, M., Carter, K.C., Ebner, R., Endress, G., Horrigan, S., Soppet, D.R. and Weaver, Z. Cancer gene determination and therapeutic screening using signature gene sets Patent: W0 0194629-A 5847 13-DEC-2001; Avalon Pharmaccuticals (US) Location/Qualifiers Corganism="Homo saplens" /db_xref="taxon:9606" T 350 a 388 c 340 g 293 t atch atch 28.4%; Score 433.2; DB 6; Length 1371;	9; Conservative
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PTENKTTPNTAEFAFSLYKOLAHQSWTSNIRFSPSVSLATAFAMLSLGFTADTHDELLE
GLURNLTEIPEAQIHEGFQELLATLAQPDSGLQLTGGNGLELSEGLKLVDKFLEDVKK
LYHSEAFTVNFGDTEEAKQINDYVEKGTQGKIVDLVKELDRDTVFALVNYIFFKGWF
ETPEFWGNTEBEDFHVQVTVVVPWARRALGAMFNIGHCKKKASSWVLLAKYLGNATAIF
FLPDGGKLQHLENEITHDITTRFLENEDRRSASLHLPKLSTGTYDLKSVLGGLGITK
VFSNGADLSGVTEEAPLKLSKAVHKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNK
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Colau, B., Chuchana, P. and Bollen, A.
Revised sequence of full-length complementary DNA coding for human
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Bollen, A., Herzog, A., Cravador, A., Herion, P., Chuchana, P., Vander Straten, A., Loriau, R., Jacobs, P. and van Elsen, A.
Cloning and expression in Escherichia coli of full-length complementary DNA coding for human alpha 1-antitrypsin DNA 2 (4), 255-264 (1983)
                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                1153 CTGGGGCCATGTTTTAGAGGCCATACCCATGTCTATCCCCCCGGAGGTCAAGTTCAACA 1212
                                                                                     1 (bases 1 to 274)

Kurachi, K., Chandra, T., Degen, S.J., White, T.T., Marchioro, T.L., Woo, S.L. and Davie, E.W.
Cloning and sequence of cDNA coding for alpha 1-antitrypsin Proc. Natl. Acad. Sci. U.S.A. 78 (11), 6826-6830 (1981)
82082539
CCGCCCCTATGTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATTAATA
                                                                 AACCATTCGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCCATTGTTTATGGGTAAGG
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/protein_id="CAA2538.1"
/db_xref="G1:28966"
/db_xref="SWISS-PROT:P01009"
                                                                                                                                                                                                                                                                              Human mRNA for alpha 1-antitrypsin.
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Nature 312 (5989), 77-80 (1984)
85036645
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Pred. No. 5.5e-83;
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AK026174 11399 bp mRNA 11near PRI 29-SEP-2000 Homo sapiens CDNA: FLJ22521 fis, clone HRC12386, highly similar to AF113676 Homo sapiens clone FLB2803 PRO0684 mRNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kawabata, A., Hikiji,T., Kobatake,N., Inagaki,H., Ikema,Y.,
Okamoto,S., Okitani,R., Ota,T., Suzuki,Y., Obayashi,M., Nishi,T.,
Shibahara,T., Tanaka,T., Nakamura,Y., Isogai,T. and Sugano,S.
NEDO human cDNA sequencing project
Unpublished
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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1258 TTACCAAAGTTTTTTTTAACGGTGCCGATTTGAGTGGTGTTACTGAAGAAGCTCCATTAA 1317
                                                                                                                                                                                                                                                    CCGGCGCCTATGCTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATA 1437
                                                                                                                                                                                                                                                                                            CTGGGGCCATGTTTTAGAGGCCATACCCATGTCTATCCCCCCGAGGTCAAGTTCAACA 1212
                                                                                                                                                                                                                                                                                                                                                                   oligo capping; fis (full insert sequence).
Homo sapiens primary human renal epithelial cells cDNA to mRNA,
clone_llb:HRC clone:HRC12386.
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Shibahara, T., Tanaka, T. and Nakamura, Y.
Direct Submission
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/cell_type="primary human renal epithelial cells"
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/note="cloning vector pME18SFL3"
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                                     ö
Length 1399;
                                     Indels
                                     0; Mismatches 493;
 DB 9;
Score 433.2; DB 9
Pred. No. 5.5e-83;
28.4%;
59.7%;
                   Best Local Similarity 59.7
Matches 729; Conservative
 Query Match
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PN JP1984091886-A/1
PD 26-MAY-1984
PF 12-MG-1983 JP 1983147907
PR 13-AG-1982 US 82 408059, 28-APR-1983 US 83 489406 PI
PR 13-AG-1982 US 82 408059, 28-APR-1983 US 83 489406 PI
PR 13-AG-1982 US 82 408059, 28-APR-1983 US 83 CONTENT ADD UTSUDOBERII
PC C1209/99,
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Mammalia; Eutheria; Primates, Catarrhini, Hominidae, Homo.
                                                                          1158 CIGGGGCCATGTTTTAGAGGCCATACCCATGTCTATCCCCCCGGGGTCAAGTTCAACA 1217
                                                                                                       AACCATTCGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCATTGTTTATGGGTAAGG 1497
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                                                CCGGCGCTATGTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATA
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IS 47..1303
IS 47..1303
IS 1304..1434
I..1434
II..1434
II..1300
IL_peptide 119..1300
ILocation/Qualifiers
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JIMOSU CORP
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strandedness: Double;
topology: Linear;
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/organism="Homo sapiens"
/db_xref="taxon:9606"
432 c 340 g 29
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E00195.1 GI:2168491
JP 1984091886-A/1.
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Guren, E.K. and Richiyaado, U.
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CAAATTTAGCCGAATTTGCTTTTTTTTTTAGACAATTAGCTCATCAAAGTAATTCTA 477
                                201 CCAACCTGGCTGAGTTCGCCTTCAGCCTATACCGCCAGCTGGCACACCAGTCCAACAGCA 260
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                                                                                       PAT 02-DEC-1994
                                                                                                                                                                                                                Method of expressing alpha-1-antitrypsin in bacteria and its use in
therapeutit formulations, and vectors and bacteria for such method
and their production
Patent: EP 0137633-Al 3 17-APR-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCAAGGCTGACACTCACGATGAAATCCTGGAGGCCTGAATTTCAACCTCACGGAGATTC 380
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                                                                                 1434 bp
Sequence 3 from Patent EP 0137633.
104196
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Parker,M.L. and Kawasaki,G.H.
                                                                                                                                                                                                                                                                   Location/Qualifiers
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432 c 340 g
TTGTCAACCCAACTCAGAAGTA 1519
                      1281 TGGTGAATCCCACCCAAAATA 1302
                                                                                                                         104196.1 GI:591838
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Russell.P.R.
Method for expressing foreign genes in schizosaccharomyces pombe and the use in therapeutic formulations of the products, DNA constructs and transformant strains of schizosaccharomyces pombe usable in such method and their preparation
patent: EP 0139383-A1 1 02-MAY-1985;
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                                             958 TTCATGTTGATCAAGTTACTACTGTCAAAGTTCCAATGATGAAAAGACTGGGTATGTTCA 1017
                                                                                                                                                  1018 ATATICAACATIGCAAAAATTAAGITCITGGGICTTATIAATGAAGTATTTAGGIAACG 1077
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Sequence 1 from Patent EP 0139383.
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Kawasaki,G.H. and Woodbury,R.G.
Glycolytic promoters for regulated protein expression: protease
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Pred. No. 5.5e-83;
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Location/Qualifiers
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107849.1 GI:589437
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